SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: McBride, William Dean, Richard T.
 - (ii) TITLE OF INVENTION: Monoamine, Diamide, Thiol-containing Metal Chelating Agents
 - (iii) NUMBER OF SEQUENCES: 9
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Banner & Allegretti, Ltd.
 - (B) STREET: 10 South Wacker Drive Suite 3000
 - (C) CITY: Chicago
 - (D) STATE: Illinois
 - (E) COUNTRY: USA
 - (F) ZIP: 60606
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/253,973
 - (B) FILING DATE: 03-JUN-1995
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Noonan, Kevin E
 - (B) REGISTRATION NUMBER: 35,303
 - (C) REFERENCE/DOCKET NUMBER: 91,875-J
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 312 715 1000
 - (B) TELEFAX: 312 715 1234
 - (C) TELEX: 910-221-5317
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 19
 - (D) OTHER INFORMATION: /label= Amide

/note= "The carboxyl terminus is modified to an
amide"

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
- Gly Gly Cys Ser Ile Pro Pro Glu Val Lys Phe Asn Lys Pro Phe Val 1 5 15

Tyr Leu Ile

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Gly Gly Cys Ser Ile Pro Pro Glu Val Lys Phe Asn Lys Pro Phe Val 1 5 10 15

Tyr Leu Ile

- (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Gly Gly Cys Gly Leu Phe 1 5

- (2) INFORMATION FOR SEQ ID NO:4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 19
 - (D) OTHER INFORMATION: /label= Amide /note= "The carboxyl terminus is modified to an amide"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Arg Gly Cys Ser Ile Pro Pro Glu Val Lys Phe Asn Lys Pro Phe Val
1 10 15

Tyr Leu Ile

- (2) INFORMATION FOR SEQ ID NO:5:
 - (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Arg Gly Cys Gln Ala Pro Leu Tyr Lys Lys Ile Ile Lys Lys Leu 1 5 10 15

Leu Glu Ser

V" 5 0

- (2) INFORMATION FOR SEQ ID NO:6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 30 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 30
 - (D) OTHER INFORMATION: /label= Amide
 /note= "The carboxyl terminus is modified to an
 amide"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Arg Gly Cys Gly His Arg Pro Leu Asp Lys Lys Arg Glu Glu Ala Pro 1 10 15

Ser Leu Arg Pro Ala Pro Pro Pro Ile Ser Gly Gly Tyr Arg
20 25 30

- (2) INFORMATION FOR SEQ ID NO:7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FRATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 14
 - (D) OTHER INFORMATION: /label= Amide /note= "The carboxyl terminus is modified to an amide"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
 - Gly Gly Cys Arg Pro Lys Pro Gln Gln Phe Phe Gly Leu Met
 1 10

(2) INFORMATION FOR SEQ ID NO:8:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 7
 - (D) OTHER INFORMATION: /label= D-Phe /note= "The phenylalanine residue is in the Dstereochemical configuration"
- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 9
 - (D) OTHER INFORMATION: /label= D-Trp /note= "The tryptophan residue is in the Dstereochemical configuration"
- (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 (B) LOCATION: 13

 - (D) OTHER INFORMATION: /label= Amide /note= "The carboxyl terminus is modified to an amide"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Ala Lys Cys Gly Gly Gly Phe Tyr Trp Lys Thr Phe Thr

- (2) INFORMATION FOR SEQ ID NO:9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 8
 - (D) OTHER INFORMATION: /label= Amide /note= "The carboxyl terminus is modified to an amide"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
 - Gly Gly Cys Phe Val Tyr Leu Ile